



#8/C

SUBSTITUTE SEQUENCE LISTING

<110> GVS Gesellschaft Fur Erwerb und Verwertung von Schutzrechten-GVS mbH and
Forschungszentrum Borstel
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Heinz, Ernst
Zahringer, Ulrich

#8/C

<120> Processive Sugar Transferase

<130> MAIWAM2.001CP1

<140> US 09/668,788

<141> 2000-09-22

<150> PCT/DE99/00857

<151> 1999-03-25

<150> DE 198 13 017.1

<151> 1998-03-25

<150> DE 198 19 958.9

<151> 1998-05-05

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<170> FastSEQ for Windows Version 4.0

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35 40 45
Ile Val Ser Glu Val Thr Gln Tyr Leu Tyr Leu Lys Ser Phe Ser Ile
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Gly Lys Gln Phe Tyr Arg Leu Phe Tyr Tyr Gly Val Asp Lys Ile Tyr
65 70 75 80
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Pro Thr Phe Asn Val Met Thr Asp Phe Cys Leu His Lys Ile Trp Val
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His Glu Asn Val Asp Lys Tyr Tyr Val Ala Thr Asp Tyr Val Lys Glu
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Lys Leu Leu Glu Ile Gly Thr His Pro Ser Asn Val Lys Ile Thr Gly
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Tyr Lys Lys Tyr Asn Leu Ser Pro Asn Lys Lys Val Leu Leu Ile Met
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210 215 220
Leu Val Lys Asp Asp Gln Val Gln Val Val Val Cys Gly Lys Asn
225 230 235 240
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245 250 255
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 35 40 45
 Ala His Pro Ile Leu Thr Ser Ile Cys Lys Lys Trp Tyr Ile Asn Ser
 50 55 60
 Phe Lys Tyr Phe Arg Asn Met Tyr Lys Gly Phe Tyr Tyr Ser Arg Pro
 65 70 75 80
 Asp Lys Leu Asp Lys Cys Phe Tyr Lys Tyr Tyr Gly Leu Asn Lys Leu
 85 90 95
 Ile Asn Leu Leu Ile Lys Glu Lys Pro Asp Leu Ile Leu Leu Thr Phe
 100 105 110
 Pro Thr Pro Val Met Ser Val Leu Thr Glu Gln Phe Asn Ile Asn Ile
 115 120 125
 Pro Val Ala Thr Val Met Thr Asp Tyr Arg Leu His Lys Asn Trp Ile
 130 135 140
 Thr Pro Tyr Ser Thr Arg Tyr Tyr Val Ala Thr Lys Glu Thr Lys Gln
 145 150 155 160
 Asp Phe Ile Asp Val Gly Ile Asp Pro Ser Thr Val Lys Val Thr Gly
 165 170 175
 Ile Pro Ile Asp Asn Lys Phe Glu Thr Pro Ile Asn Gln Lys Gln Trp
 180 185 190
 Leu Ile Asp Asn Asn Leu Asp Pro Asp Lys Gln Thr Ile Leu Met Ser
 195 200 205

Ala Gly Ala Phe Gly Val Ser Lys Gly Phe Asp Thr Met Ile Thr Asp
 210 215 220
 Ile Leu Ala Lys Ser Ala Asn Ala Gln Val Val Met Ile Cys Gly Lys
 225 230 235 240
 Ser Lys Glu Leu Lys Arg Ser Leu Thr Ala Lys Phe Lys Leu Thr Arg
 245 250 255
 Met Tyr Leu Ile Leu Gly Tyr Thr Lys His Met Asn Glu Trp Met Ala
 260 265 270
 Ser Ser Gln Leu Met Ile Thr Lys Pro Gly Gly Ile Thr Ile Thr Glu
 275 280 285
 Gly Phe Ala Arg Cys Ile Pro Met Ile Phe Leu Asn Pro Ala Pro Gly
 290 295 300
 Gln Glu Leu Glu Asn Ala Phe Tyr Phe Glu Glu Lys Gly Phe Gly Lys
 305 310 315 320
 Ile Ala Asp Thr Pro Glu Glu Ala Ile Lys Ile Val Ala Ser Leu Thr
 325 330 335
 Asn Gly Asn Glu Gln Leu Thr Asn Met Ile Ser Thr Met Glu Gln Asp
 340 345 350
 Lys Ile Lys Tyr Ala Thr Gln Thr Ile Cys Arg Asp Leu Leu Asp Leu
 355 360 365
 Ile Gly His Ser Ser Gln Pro Gln Glu Ile Tyr Gly Lys Val Pro Leu
 370 375 380
 Tyr Ala Arg Phe Phe Val Lys
 385 390